Complexity Measures: Linguistic and Symbolic Dynamical Approaches



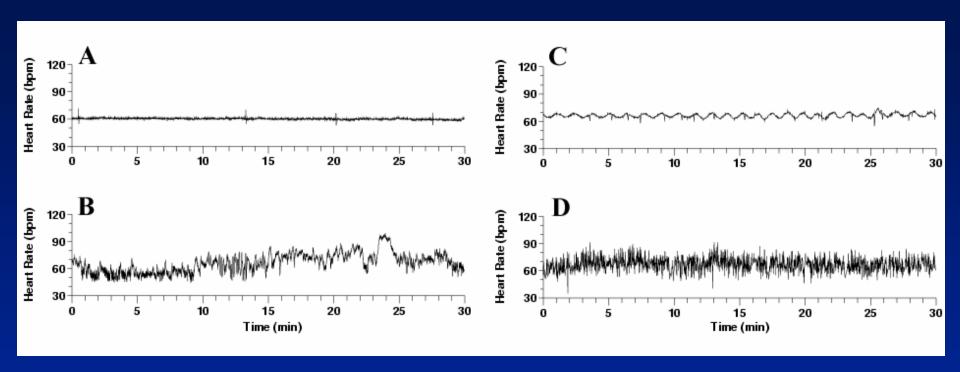
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Objectives

- Provide an analogy between heart rate time series and other symbolic sequences (e.g., language)
- Introduce an intuitive measurement of similarity based on repetitive patterns
- Discuss clinical applications of this linguistic approach

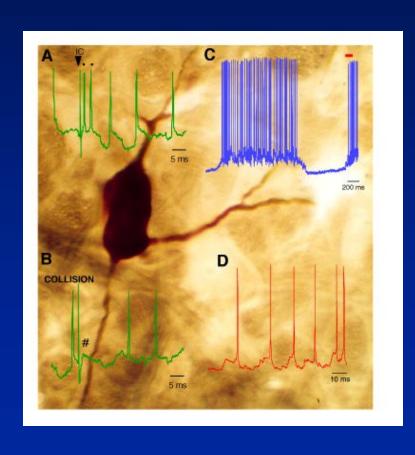
Heart Rate Dynamics in Health and Disease

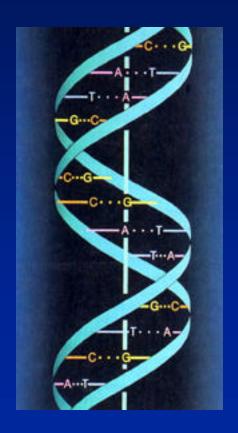


These time series contain information that reflects their underlying control mechanisms

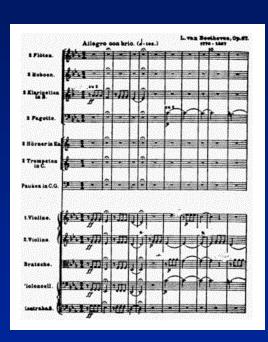
Question: Can we usefully classify different types of information even *without* initially understanding their content?

Information Created by Biological Systems

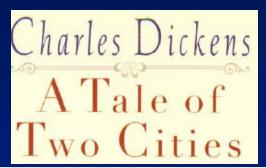




Information Created by Humans







1 The Period

IT WAS the best of times, it was the worst of times, it was the age of wisdom, it was the age of foolishness, it was the epoch of belief, it was the epoch of incredulity, it was the season of Light, it was the season of Darkness, it was the spring-of hope, it was the winter of despair, we had everything before us, we had nothing before us, we were all going direct to Heaven, we were all going direct the other way—in short, the period was so far like the present period, that some of its noisiest authorities insisted on its being received, for good or for evil, in the superlative degree of comparison only.

Our approach: Categorize complex signals based on the occurrences of *repetitive patterns*

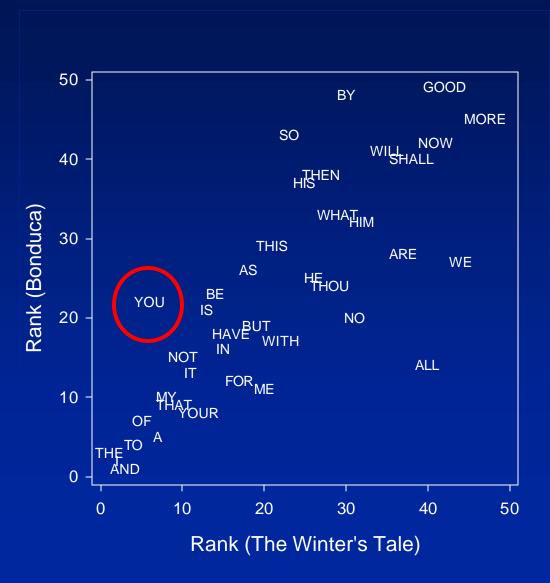
Illustration of our algorithm: Comparison of human literary texts

Repetitive patterns: words

Refs: Phys Rev Lett 2003; 90:108103; Physica A 2003:329:473-83

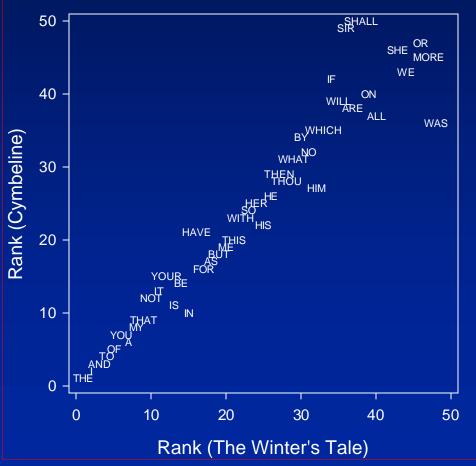
Comparison of Human Texts

Word	Rank (The Winter's Tale)	Rank (Bonduca)
THE	1	3
I	2	2
AND	3	1
TO	4	4
OF	5	7
YOU	6	22
A	7	5
MY	8	10
THAT	9	9
NOT	10	15

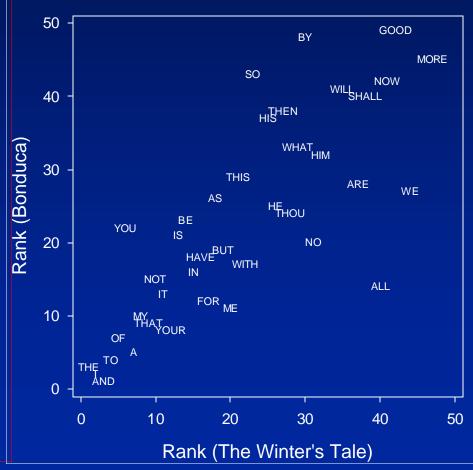


Rank Comparison Maps

Shakespeare vs. Shakespeare



Shakespeare vs. Fletcher

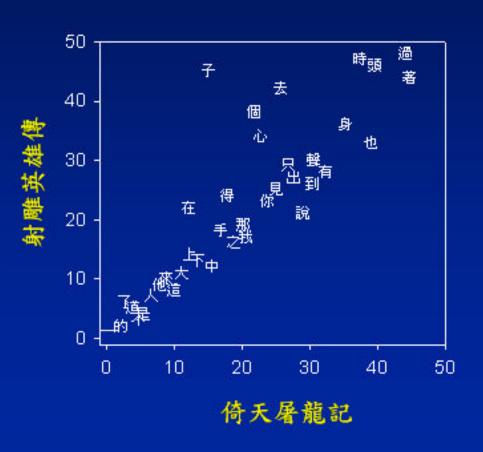


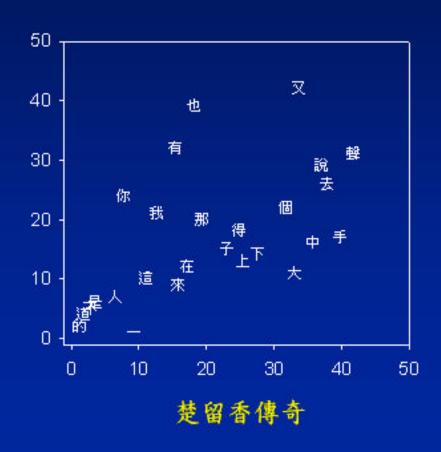


Rank Comparison Maps

Same Author

Different Authors

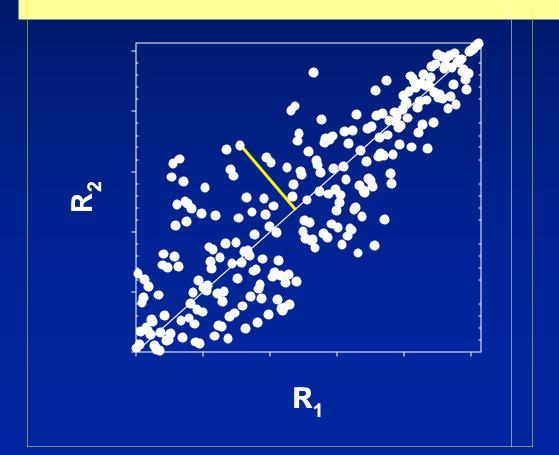




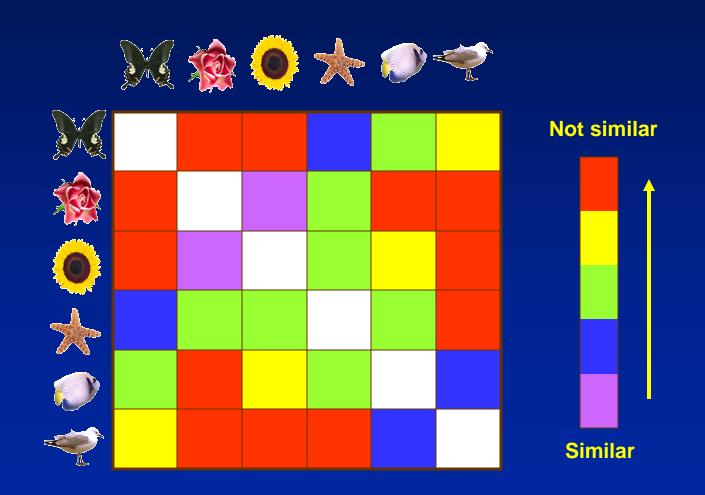
Dis-Similarity Index

$$D(S_1, S_2) = \sum_{k=1}^{N} |R_1(w_k) - R_2(w_k)| F(w_k)$$

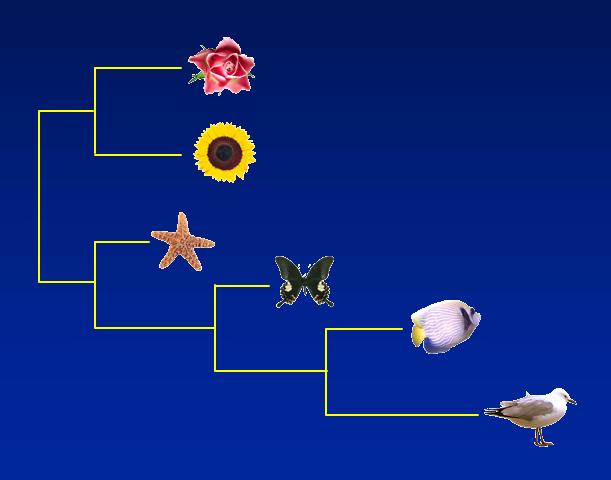
$$F(w_k) = \frac{1}{Z} \left[-p_1(w_k) \log p_1(w_k) - p_2(w_k) \log p_2(w_k) \right]$$



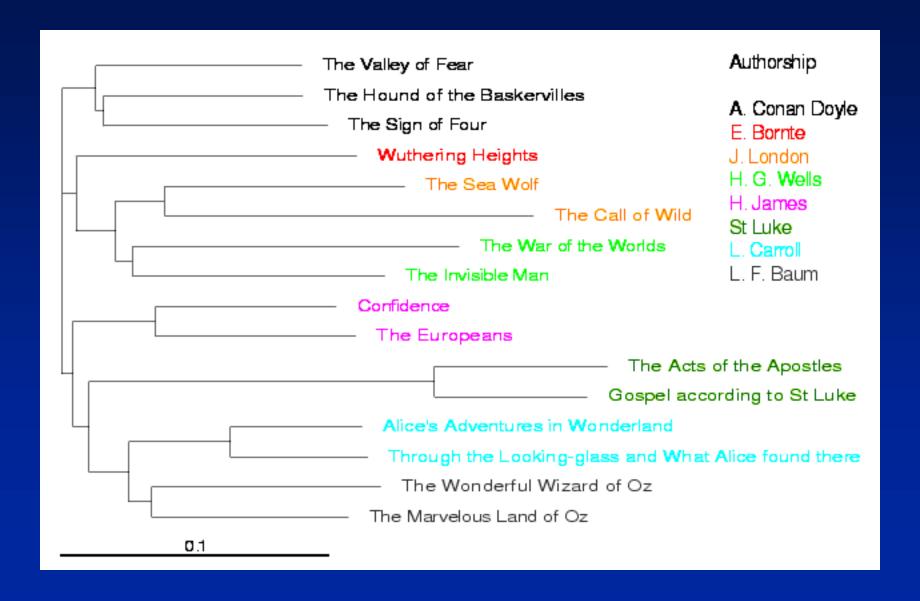
Phylogenetic Tree Distance matrix method



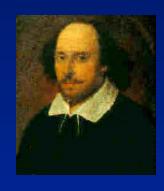
Phylogenetic Tree



Authorship Problem



Who Wrote Shakespeare's Plays?



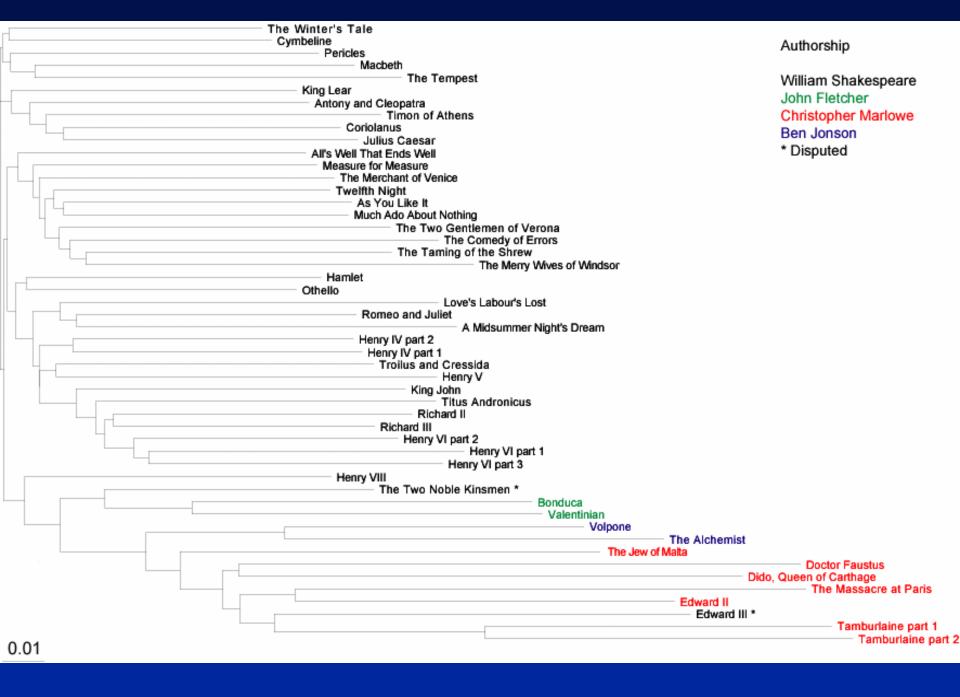


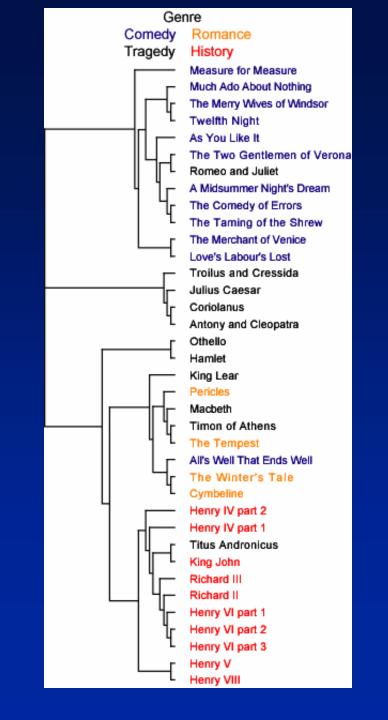


Shakespeare

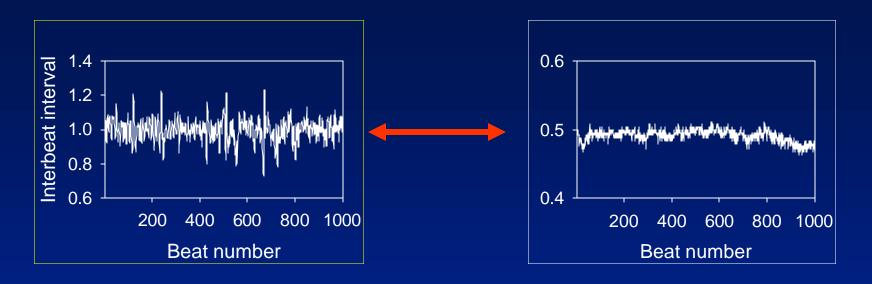


Marlowe



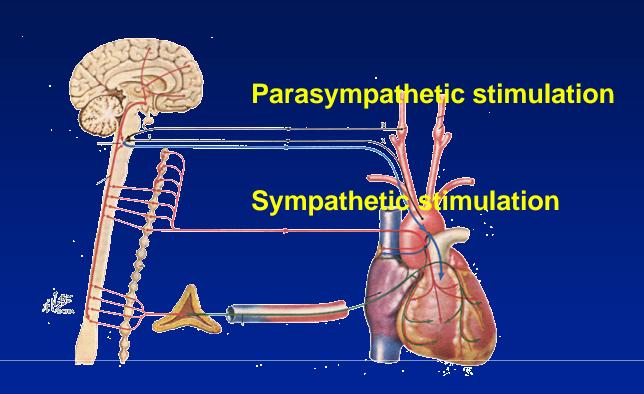


Application to Heartbeat Time Series



- How to map a time series to a symbolic sequence?
- How to define words in symbolic sequences?

Heart rate dynamics





Symbolic Mapping

Step 1: Binary code representation

Increment of successive RR intervals can be mapped to 0 or 1

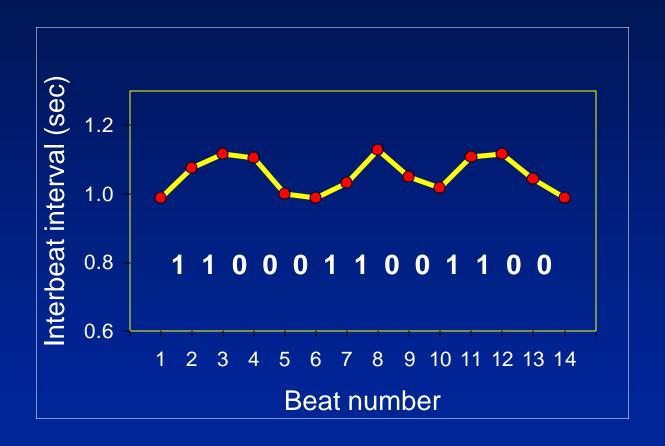
$$I_n = \begin{cases} 0, & \text{if } x_n \le x_{n-1} \\ 1, & \text{if } x_n > x_{n-1} \end{cases}$$

Ashkenazy et al., Phys. Rev. Lett. 86, 1900 (2001).

Step 2: "Word" partitioning

m+1 successive RR intervals are mapped to a binary sequence of length m, called an **m-bit word**

Symbolic Mapping



8-bit word: 11000110, 10001100, 00011001

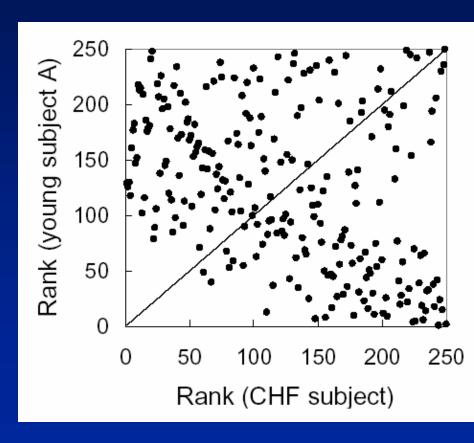
Scientific Questions

• As a healthy system is altered by disease and aging, can we quantify changes in the dynamical patterns?

Health vs. Health

Rank (young subject A) Rank (young subject B)

Health vs. Disease



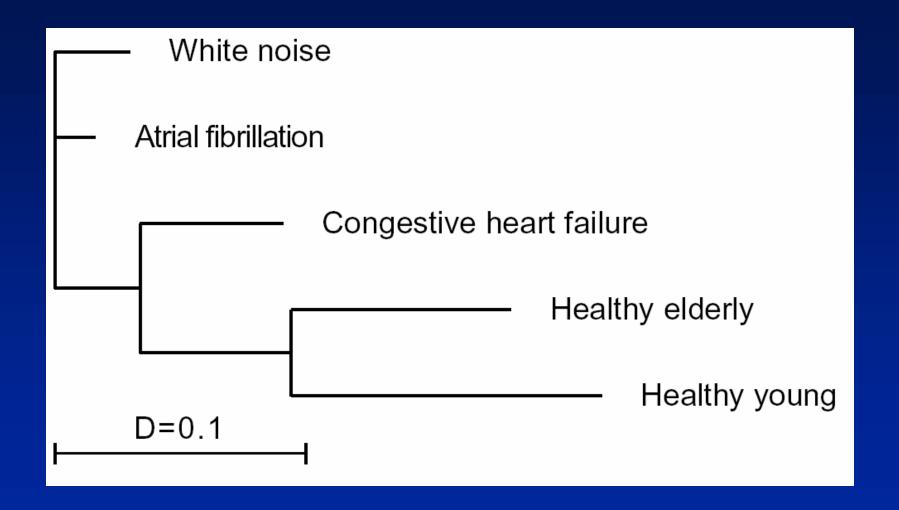
D = 0.10

D = 0.25

PhysioNet: www.physionet.org NIH Research Resource for Complex Physiologic Signals

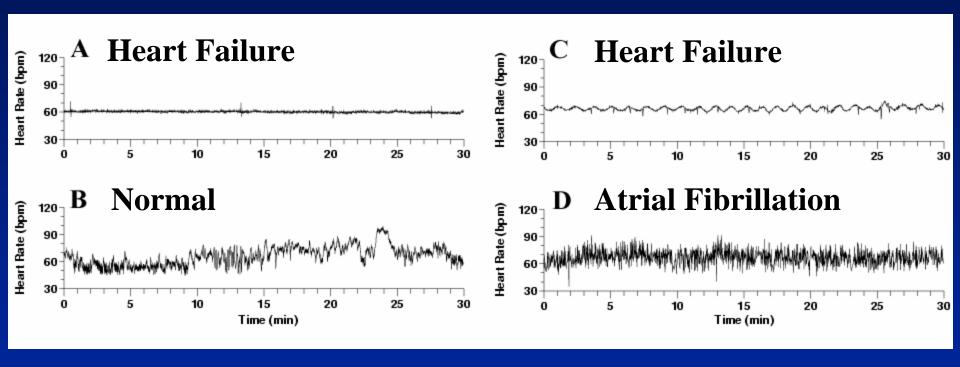
- Healthy subjects 2 hours
 Young (10 male, 10 female, average 25.9 years)
 Elderly (10 male, 10 female, average 74.5 years)
- Congestive heart failure database 16-24 hours
 15 female, 28 males, average 55.5 years
- Atrial fibrillation database 2 hours9 subjects
- White noise (random Gaussian noise)

Phylogenetic Tree



Yang et al. Phys Rev Lett, 2003; 90: 108103

Heart Rate Dynamics in Health and Disease

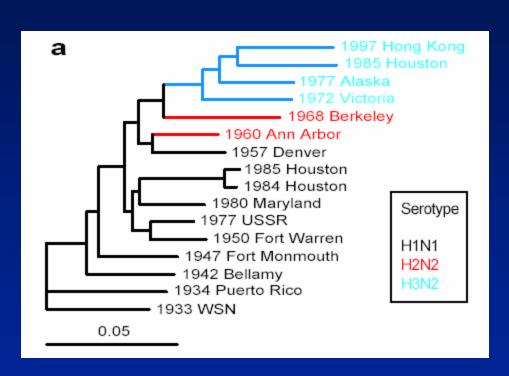


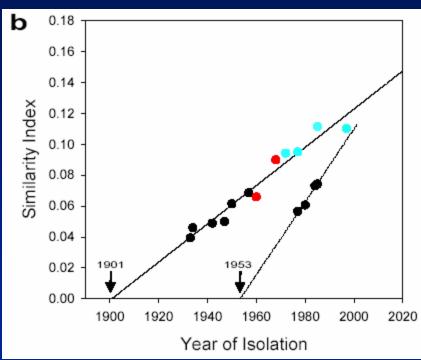
It is possible to categorize these time series!

Other Applications

- Gene chip (micro-array) analysis
- Dynamics of sleep stage transitions
- EEG classification
- Image analysis (pathology)
- Psychiatric evaluation

Application to DNA Sequences: Human Influenza Virus



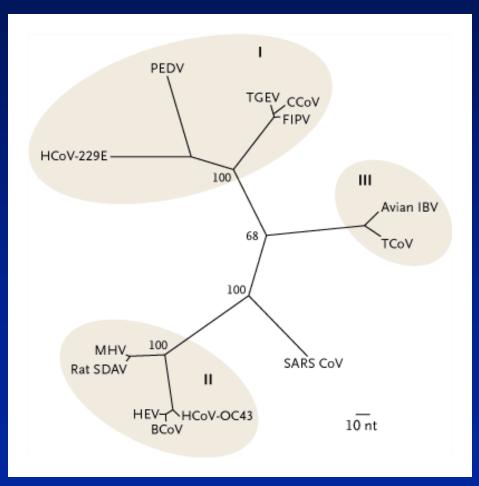


Our result is consistent with previous finding based on sequence alignment technique (*Science* 1986; **232**: 980)

Classification of SARS Virus (~29751 bps)

WHO Result (based on **405 bps**)

Our result (complete genome)



SARS-CoV **TGEV** HCoV-229E **PEDV BCoV IBV** MHV D=0.05

N Engl J Med 2003; **348**: 1947

Yang, Goldberger, Peng. J Comput Biol 2005;12:1103-1116

Conclusions

- Biological signals are complex and contain hidden information
- Information categorization analysis based on repetitive elements may uncover certain interesting features relevant to modeling and diagnostics
- Applications range from literary texts, DNA sequences, to physiologic time series